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RMS Patent Department

NO. 125 P. 6

Application No. 09/823,649

Appendix B
(4 pages)

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 European Bioinformatics Institute Get Nucleotide sequences for <input type="text"/> <input type="button" value="Get?"/> Site search <input type="text"/> <input type="button" value="Go?"/>		<input type="button" value="Home"/> <input type="button" value="Site Map"/> <input type="button" value="Database Queries"/>																	
<table border="1"> <tr> <td>InterPro home</td> <td>Text Search</td> <td>Sequence Search</td> <td>Databases</td> </tr> <tr> <td colspan="2"><input type="text"/> Search:</td> <td><input type="button" value="Search Entries"/></td> <td><input type="button" value="Documentation"/></td> </tr> <tr> <td colspan="2"></td> <td><input type="button" value="FTP site"/></td> <td><input type="button" value="Protein of the month"/></td> </tr> <tr> <td colspan="4"> Simple Full HTML Version Click here for help! </td> </tr> </table>				InterPro home	Text Search	Sequence Search	Databases	<input type="text"/> Search:		<input type="button" value="Search Entries"/>	<input type="button" value="Documentation"/>			<input type="button" value="FTP site"/>	<input type="button" value="Protein of the month"/>	Simple Full HTML Version Click here for help!			
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<h3>InterPro Protein matches</h3> <table border="1"> <tr> <td>Select: <input type="radio"/></td> <td>Protein accession(s): <input type="text" value="P00582"/></td> </tr> <tr> <td>Refine: <input type="radio"/></td> <td>Proteins with alternative splicing: <input type="checkbox"/></td> </tr> <tr> <td>Display: <input type="radio"/></td> <td>Proteins with known structure: <input type="checkbox"/></td> </tr> <tr> <td>Sort: <input type="radio"/></td> <td>Taxonomy - NCBI tax ID(s): <input type="text"/></td> </tr> <tr> <td></td> <td><input type="button" value="Detailed"/></td> </tr> <tr> <td></td> <td><input type="button" value="Accession"/></td> </tr> <tr> <td colspan="2"> <input type="button" value="Match data"/>  </td> </tr> </table>				Select: <input type="radio"/>	Protein accession(s): <input type="text" value="P00582"/>	Refine: <input type="radio"/>	Proteins with alternative splicing: <input type="checkbox"/>	Display: <input type="radio"/>	Proteins with known structure: <input type="checkbox"/>	Sort: <input type="radio"/>	Taxonomy - NCBI tax ID(s): <input type="text"/>		<input type="button" value="Detailed"/>		<input type="button" value="Accession"/>	<input type="button" value="Match data"/> 			
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<h3>Taxonomy Lineage</h3> <p>no rank: Root node of taxonomy 1</p> <p>no rank: cellular organisms 1</p>																			
<table border="1"> <tr> <td>Protein </td> <td>Match data </td> </tr> <tr> <td colspan="2">DNA polymerase I (EC 2.7.7.7) (POL I)</td> </tr> <tr> <td colspan="2"> <ul style="list-style-type: none"> Table of Matches GO annotation View protein UniProt information </td> </tr> </table>				Protein 	Match data 	DNA polymerase I (EC 2.7.7.7) (POL I)		<ul style="list-style-type: none"> Table of Matches GO annotation View protein UniProt information 											
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IPR003583	SM00278	[...]
IPR008918	SM00279	[...]
IPR008918	SSF47807	[...]
IPR012337	SSFS3098	[...]
Structural features [1]		
1dp1	1dp1	[...]
1.10.150.20.17	2kfnA4	[...]
1.2D.1060.10.1	2kfnA2	[...]
3.30.420.10.16	2kfnA1	[...]
3.30.70.370.1	2kfnA3	[...]
c.55.3.5	d2fmna1	[...]
e.8.1.1	d2fmna2	[...]
Structural predictions [2]		
MB_P00582		

ModBase	[...]
PRINTS	[...]
Pfam	[...]
PROSITE pattern	[...]
SMART	[...]
SuperFamily	[...]
PDB Chain	[...]

Match Status key: True

CATH Domain	[REDACTED]
TIGRFAMs	[REDACTED]
SCOP Domain	[REDACTED]
IPR001098 DNA-directed DNA polymerase	DNA-directed DNA polymerase
IPR002421 5'-3' exonuclease	5'-3' exonuclease
IPR002562 3'-5' exonuclease	3'-5' exonuclease

F	False
P	Partial
U	Unknown
N	False negative

InterPro 12.1

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